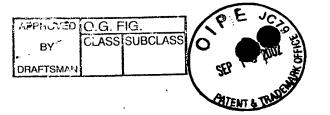


EcoRI RBS PelB leader 1 GAATTCATTAAA<u>GAGGAG</u>AAATTAACC**ATG**AAATACCTATTGCCTACGGCAGCCGCTGGCTTGCTG 1 M K Y L L P T A A A G L L Ncol ◆ VH anti-CD3 Frame-H1 67 CTGCTGGCAGCTCAGCCGGCCATGGCGCAGGTGCAGCTGCAGCAGTCTGGGGGCTGAACTGGCAAGAC 14 L L A A Q P A M A Q V Q L Q Q S G A E L A R 134 CTGGGGCCTCAGTGAAGATGTCCTGCAAGGCTTCTGGCTACACCTTTACTAGGTACACGATGCA 36 P G A S V K M S C K A S G Y T F T R Y Frame-H2 CDR-H2 198 CTGGGTAAAACAGAGGCCTGGACAGGGTCTGGAATGGATTGGATAATCCTAGCCGTGG 57 W V K Q R P G Q G L E W I G Y I N P S Frame-H3 261 TTATACTAATTACAATCAGAAGTTCAAGGACAAGGCCACATTGACTACAGACAAATCCTCCA Y T N Y N Q K F K D K A T L T T D K S S 323 GCACAGCCTACATGCAACTGAGCAGCCTGACATCTGAGGACTCTGCAGTCTATTACTGTGCAAGA<u>T</u>A 99 S T A Y M Q L S S L T S E D S A V Y Y C A R Y CDR-H3 Frame-H4 390 TTATGATGATCATTACAGCCTTGACTACTGGGGCCCAAGGCCACCACTCTCACAGTCTCCTCAG Y D D H Y S LDYWGQGTTLTVSS CH1 Linker VL anti-CD19 452 CCAAAACAACACCCAAGCTTGGCGGTGATATCTTGCTCACCCAAACTCCAGCTTCTTTGGCTGTG 142 A K T T P K L G G D I L L T Q T P A S L A V CDR-L1 164 S L G Q R A T I S C K A S Q S V D Frame-L2 579 <u>TAGTTATTTGAAC</u>TGGTACCAACAGATTCCAGGACAGCCACCCAAACTCCTCATCTAT<u>GATGCA</u> 184 S Y L N W Y Q Q I P G Q P P K L L I Y D A CDR-L2 Frame-L3 643 TCCAATCTAGTTTCTGGGATCCCACCCAGGTTTAGTGGCAGTGGGTCTGGGACAGACTTCACCC 206 S N L V S G I P P R F S G S G T D F T CDR-L3 707 TCAACATCCATCCTGTGGAGAAGGTGGATGCTGCAACCTATCACTGTCAGCAAAGTACTGAGGA 227 L N I H P V E K V D A A T Y H C Frame-L4 C kappa Noti 771 <u>T</u>CCGTGGACGTTCGGTGGAGGCACCAAGCTGGAAATCAAA<u>CGGGCTGATGCT</u>GCGGCCGCTGGATCC 248 PWTFGGGTKLEIKRADAAAGS c-myc epitope His6 tail 838 GAACAAAAGCTGATCTCAGAAGAAGACCTAAACTCACCATCACCATCACCATCACTAAAGAT 271 E Q K L I S E E D L N S H H H H H H . 899 CT





Balll **RBS** Pel B leader 1 AGATCTATTAAAGAGGAGAAATTAACCATGAAATACCTATTGCCTACGGCAGCCGCTGGCTTGC 1 M K Y L L P T A A A G L Ncol ◆ VH anti-CD19 65 TGCTGCTGGCAGCTCAGCCGGCCATGGCGCAGGTGCAGCTGCAGCAGTCTGGGGCTGAGCTGGT 13 L L L A A Q P A M A Q V Q L Q Q S G A E L V 129 GAGGCCTGGGTCCTCAGTGAAGATTTCCTGCAAGGCTTCTGGCTATGCATTCAGT<u>AGCTACTG</u> 34 R P G S S V K I S C K A S G Y A F S S Frame-H2 192 GATGAACTGGGTGAAGCAGAGGCCTGGACAGGGTCTTGAGTGGATTGGACAGATTTGGCCT 55 M N W V K Q R P G Q G L E W I G Q I CDR-H2 253 GGAGATGGTGATACTAACTACAATGGAAAGTTCAAGGGTAAAGCCACTCTGACTGCA 76 G D G D T N Y N G K F KGKATLTA Frame-H3 310 GACGAATCCTCCAGCACAGCCTACATGCAACTCAGCAGCCTAGCATCTGAGGACTCTGCGGTCT 95 D E S S S T A Y M Q L S S L A S E D S A V CDR-H3 374 ATTTCTGTGCAAGACGGGAGACTACGACGGTAGGCCGTTATTACTATGCTATGGACT 116 Y F C A R R E T T T V G R Y Y Y A M D Frame-H4 CH₁ Linker 431 <u>AC</u>TGGGGTCAAGGAACCTCAGTCACCGTCTCCTCAGCCAAAACAACACCCCAAGCTTGGCGGT 135 Y W G Q G T S V T V S S A K T T P K L G G VL anti-CD3 Frame-L1 493 GATATCGTGCTCACTCAGTCTCCAGCAATCATGTCTGCATCTCCAGGGGAGAAGGTCACCATGA 156 D I V L T Q S P A I M S A S P G E K V T M CDR-L1 Frame-L2 557 CCTGC<u>AGTGCCAGCTCAAGTGTAAGTTACATGAAC</u>TGGTACCAGCAGAAGTCAGGCACC 177 T C S S S V S Y M N W Y Q Q K S G T CDR-L2 616 TCCCCCAAAAGATGGATTTATGACACATCCAAACTGGCTTCTGGAGTCCCTGCTCACTTC 197 S P K R W I Y D T S K L Α SGVPAHF Frame-L3 676 AGGGGCAGTGGGTCTGGGACCTCTTACTCTCTCACAATCAGCGGCATGGAGGCTGAAGATGCTG 217 R G S G S G T S Y S L T I S G M E A E D A CDR-L3 Frame-L4 740 CCACTTATTACTGCCCAGCAGTGGAGTAGTAACCCATTCACGTTCGGCTCGGGGACAAAG 238 A T Y Y C Q Q W S S N P F T F G S G T K C kappa c-myc epitope 799 TTGGAAATAAAC<u>CGGGCTGATACTGCACCAACT</u>GGATCC*GAACAAAAGCTGATCTCAGAA* 258 LEINRADTAPTGSEQKLISE His6 tail 859 GAAGACCTAAACTCACCATCACCATCACCATCACTAATCTAGA D L N S H H H H H .